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#12
KAY
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FEDERAL BUREAU OF INVESTIGATION
U.S. DEPARTMENT OF JUSTICE
1600/2900

AUG 13 2002

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/516,444A

DATE: 08/08/2002

TIME: 12:03:45

Input Set : A:\5995USP2.txt

Output Set: N:\CRF4\08082002\I516444A.raw

4 <110> APPLICANT: Abbott Laboratories
5 Billing-Medel, Patricia A.
6 Cohen, Maurice
7 Colpitts, Tracey L.
8 Friedman, Paula N.
9 Gordon, Julian
10 Granados, Edward N.
11 Hodges, Steven C.
12 Klass, Michael R.
13 Kratochvil, Jon D.
14 Roberts-Rapp, Lisa
15 Russell, John C.
16 Stroupe, Stephen D.
18 <120> TITLE OF INVENTION: Reagents And Method Useful For Detecting
19 Diseases Of The Breast
22 <130> FILE REFERENCE: 5995.US.P2
24 <140> CURRENT APPLICATION NUMBER: 09/516,444A
25 <141> CURRENT FILING DATE: 2000-02-29
27 <150> PRIOR APPLICATION NUMBER: US 08/962,094
28 <151> PRIOR FILING DATE: 1997-10-31
30 <150> PRIOR APPLICATION NUMBER: US 08/742,067
31 <151> PRIOR FILING DATE: 1996-10-31
33 <160> NUMBER OF SEQ ID NOS: 39
35 <170> SOFTWARE: FastSEQ for Windows Version 4.0
37 <210> SEQ ID NO: 1
38 <211> LENGTH: 201
39 <212> TYPE: DNA
40 <213> ORGANISM: Artificial Sequence
42 <220> FEATURE:
43 <223> OTHER INFORMATION: EST Clone 1662885
W--> 45 <221> NAME/KEY: misc_feature
46 <222> LOCATION: (26)...(26)
47 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
48 position 26
W--> 50 <221> misc_feature
51 <222> LOCATION: (98)...(98)
52 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
53 position 98
W--> 55 <221> misc_feature
56 <222> LOCATION: (133)...(133)
57 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
58 position 133
W--> 60 <221> misc_feature

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61 <222> LOCATION: (145)...(145)
62 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
63     position 145
W--> 65 <221> misc_feature
66 <222> LOCATION: (183)...(183)
67 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
68     position 183
W--> 70 <400> 1
W--> 71 ctcttaggct ttgaagcatt tttgtntgtg ctccctgac ttcatgtcac caccatgaag 60
W--> 72 ttcttagcag tcctggtagt cttgggagtt tccatctntc tggctctctgc ccagaatccg 120
W--> 73 acaacagctg ctncagctga cacgnatcca gctactggtc ctgctgatga tgaagcccct 180
W--> 74 gangctgaaa ccactgctgc t                                     201
76 <210> SEQ ID NO: 2
77 <211> LENGTH: 308
78 <212> TYPE: DNA
79 <213> ORGANISM: Homo sapiens
81 <400> SEQUENCE: 2
82 taggctttga agcatttttg tctgtgctcc ctgatottca ggtaaccacc atgaagttct 60
83 tagcagtcct ggtactcttg ggagtttcca tctttctggt ctctgccag aatccgacaa 120
84 cagctgctcc agctgacacg tatccagcta ctggtcctgc tgatgatgaa gcccctgatg 180
85 ctgaaaccac tgctgctgca accactgcga ccactgctgc tcctaccact gcaaccaccg 240
86 ctgcttctac cactgctcgt aaagacattc cagttttacc caaatgggtt ggggatcttc 300
87 cgaatggt                                     308
89 <210> SEQ ID NO: 3
90 <211> LENGTH: 292
91 <212> TYPE: DNA
92 <213> ORGANISM: Artificial Sequence
94 <220> FEATURE:
95 <223> OTHER INFORMATION: EST Clone 901429
W--> 97 <221> NAME/KEY: misc_feature
98 <222> LOCATION: (236)...(236)
99 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
100     position 236
W--> 102 <221> misc_feature
103 <222> LOCATION: (259)...(259)
104 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
105     position 259
W--> 107 <400> 3
108 gcatttttgt ctgtgctccc tgatcttcat gtcaccacca tgaagttctt agcagtcctg 60
109 gtactcttgg gagtttccat ctttctgggc tctgccaga atccgacaa agctgctcca 120
110 gctgacacgt atccagctac tggctcctgct gatgatgaag cccctgatgc tgaaaccact 180
W--> 111 gctgctgcaa ccactgcgac cactgctgct cctaccactg caaccaccgc tgcttntacc 240
W--> 112 actgctcgta aagacattnc agttttaccc aaatggggtg gggatctccc ga       292
114 <210> SEQ ID NO: 4
115 <211> LENGTH: 197
116 <212> TYPE: DNA
117 <213> ORGANISM: Homo sapiens
119 <400> SEQUENCE: 4
120 gttttaccca aatggggttg ggatctcccg aatggtagag tgtgtccctg agatggaatc 60

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121 agcttgagtc ttctgcaatt ggtcacaaact attcatgctt cctgtgattt catccaacta 120
122 cttaccttgc ctacgatata ccctttatct ctaatcagtt tattttcttt caaataaaaa 180
123 ataactatga gcaacat 197
125 <210> SEQ ID NO: 5
126 <211> LENGTH: 472
127 <212> TYPE: DNA
128 <213> ORGANISM: Homo sapiens
130 <400> SEQUENCE: 5
131 ctcttaggct ttgaagcatt tttgtctgtg ctccctgata ttcattgtcac caccatgaag 60
132 ttcttagcag tcttggtact cttgggagtt tccatctttc tggctctctgc ccagaatccg 120
133 acaacagctg ctccagctga cactgatcca gctactggtc ctgctgatga tgaagccct 180
134 gatgctgaaa ccaactgctgc tgcaaccact gcgaccactg ctgctcctac cactgcaacc 240
135 accgctgctt ctaccactgc tcgtaaagac attccagttt taccctaatg ggttggggat 300
136 ctcccgatg gtagagtgtg tccctgagat ggaatcagct tgagtcttct gcaattgggc 360
137 acaactattc atgcttctctg tgatttcatc caactactta ccttgcttac gatatccct 420
138 ttatctctaa tcagtttatt ttctttcaaa taaaaaataa ctatgagcaa ca 472
140 <210> SEQ ID NO: 6
141 <211> LENGTH: 473
142 <212> TYPE: DNA
143 <213> ORGANISM: Homo sapiens
145 <400> SEQUENCE: 6
146 ctcttaggct ttgaagcatt tttgtctgtg ctccctgata ttcattgtcac caccatgaag 60
147 ttcttagcag tcttggtact cttgggagtt tccatctttc tggctctctgc ccagaatccg 120
148 acaacagctg ctccagctga cactgatcca gctactggtc ctgctgatga tgaagccct 180
149 gatgctgaaa ccaactgctgc tgcaaccact gcgaccactg ctgctcctac cactgcaacc 240
150 accgctgctt ctaccactgc tcgtaaagac attccagttt taccctaatg ggttggggat 300
151 ctcccgatg gtagagtgtg tccctgagat ggaatcagct tgagtcttct gcaattgggc 360
152 acaactattc atgcttctctg tgatttcatc caactactta ccttgcttac gatatccct 420
153 ttatctctaa tcagtttatt ttctttcaaa taaaaaataa ctatgagcaa cat 473
155 <210> SEQ ID NO: 7
156 <211> LENGTH: 68
157 <212> TYPE: DNA
158 <213> ORGANISM: Artificial Sequence
160 <220> FEATURE:
161 <223> OTHER INFORMATION: Restriction site
163 <400> SEQUENCE: 7
164 agctcggaat tccgagcttg gatcctctag agcgggccgcc gactagttag ctcgtcgacc 60
165 cgggaatt 68
167 <210> SEQ ID NO: 8
168 <211> LENGTH: 68
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
172 <220> FEATURE:
173 <223> OTHER INFORMATION: Restriction site
175 <400> SEQUENCE: 8
176 aattaattcc cgggtcgacg agctcactag tcggcgggccg ctctagagga tccaagctcg 60
177 gaattccg 68
179 <210> SEQ ID NO: 9
180 <211> LENGTH: 24

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181 <212> TYPE: DNA
182 <213> ORGANISM: Artificial Sequence
184 <220> FEATURE:
185 <223> OTHER INFORMATION: Universal Primer
187 <400> SEQUENCE: 9
188 agcggataac aatttcacac agga                24
190 <210> SEQ ID NO: 10
191 <211> LENGTH: 18
192 <212> TYPE: DNA
193 <213> ORGANISM: Artificial Sequence
195 <220> FEATURE:
196 <223> OTHER INFORMATION: Universal Primer
198 <400> SEQUENCE: 10
199 tgtaaaacga cggccagt                18
201 <210> SEQ ID NO: 11
202 <211> LENGTH: 20
203 <212> TYPE: DNA
204 <213> ORGANISM: Artificial Sequence
206 <220> FEATURE:
207 <223> OTHER INFORMATION: Primer
209 <400> SEQUENCE: 11
210 actgctcgta aagacattcc                20
212 <210> SEQ ID NO: 12
213 <211> LENGTH: 19
214 <212> TYPE: DNA
215 <213> ORGANISM: Artificial Sequence
217 <220> FEATURE:
218 <223> OTHER INFORMATION: Primer
220 <400> SEQUENCE: 12
221 gggacacact ctaccattc                19
223 <210> SEQ ID NO: 13
224 <211> LENGTH: 20
225 <212> TYPE: DNA
226 <213> ORGANISM: Artificial Sequence
228 <220> FEATURE:
229 <223> OTHER INFORMATION: Sense Primer
231 <400> SEQUENCE: 13
232 aagcccctga tgctgaaacc                20
234 <210> SEQ ID NO: 14
235 <211> LENGTH: 23
236 <212> TYPE: DNA
237 <213> ORGANISM: Artificial Sequence
239 <220> FEATURE:
240 <223> OTHER INFORMATION: Antisense Primer
242 <400> SEQUENCE: 14
243 tgcagaagac tcaagctgat tcc                23
245 <210> SEQ ID NO: 15
246 <211> LENGTH: 20
247 <212> TYPE: DNA

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248 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: Target-Specific Forward Primer
253 <400> SEQUENCE: 15
254 aagccctga tgctgaaacc                                20
256 <210> SEQ ID NO: 16
257 <211> LENGTH: 23
258 <212> TYPE: DNA
259 <213> ORGANISM: Artificial Sequence
261 <220> FEATURE:
262 <223> OTHER INFORMATION: Target-Specific Reverse Primer
264 <400> SEQUENCE: 16
265 tgcagaagac tcaagctgat tcc                            23
267 <210> SEQ ID NO: 17
268 <211> LENGTH: 15
269 <212> TYPE: DNA
270 <213> ORGANISM: Artificial Sequence
272 <220> FEATURE:
273 <223> OTHER INFORMATION: Probe
275 <400> SEQUENCE: 17
276 gaccactgct gctcc                                    15
278 <210> SEQ ID NO: 18
279 <211> LENGTH: 20
280 <212> TYPE: DNA
281 <213> ORGANISM: Artificial Sequence
283 <220> FEATURE:
284 <223> OTHER INFORMATION: Sense Primer
286 <400> SEQUENCE: 18
287 actgctcgta aagacattcc                                20
289 <210> SEQ ID NO: 19
290 <211> LENGTH: 19
291 <212> TYPE: DNA
292 <213> ORGANISM: Artificial Sequence
294 <220> FEATURE:
295 <223> OTHER INFORMATION: Antisense Primer
297 <400> SEQUENCE: 19
298 gggacacact ctaccattc                                19
300 <210> SEQ ID NO: 20
301 <211> LENGTH: 90
302 <212> TYPE: PRT
303 <213> ORGANISM: Homo sapiens
305 <400> SEQUENCE: 20
306 Met Lys Phe Leu Ala Val Leu Val Leu Leu Gly Val Ser Ile Phe Leu
307 1          5          10          15
308 Val Ser Ala Gln Asn Pro Thr Thr Ala Ala Pro Ala Asp Thr Tyr Pro
309 20          25          30
310 Ala Thr Gly Pro Ala Asp Asp Glu Ala Pro Asp Ala Glu Thr Thr Ala
311 35          40          45
312 Ala Ala Thr Thr Ala Thr Thr Ala Ala Pro Thr Thr Ala Thr Thr Ala

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VERIFICATION SUMMARY

DATE: 08/08/2002

PATENT APPLICATION: US/09/516,444A

TIME: 12:03:46

Input Set : A:\5995USP2.txt

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L:45 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:50 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:60 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:65 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:70 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:71 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:60
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:120
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180
L:97 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:102 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:107 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:180
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:240



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Team: OIPEBackFileIndexing

Dossier: 09516444

Legal Date: 10-02-2003

No.	Doccode	Number of pages
1	CTRS	6
2	NFDR	1
3	EXIN	1

Total number of pages: 8

Remarks:

Order of re-scan issued on